

GenCore version 5.1.6  
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OM protein - protein search, using sw model

Run on: December 3, 2005, 23:21:32 ; Search time 5 Seconds  
(without alignments)  
261.443 Million cell updates/sec

Perfect score: 1505  
Sequence: 1 MRGSQVLLMWLEVLAVGGT.....SEQISFLEQLGSCSKCKDS 273

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 26661 seqs, 4788334 residues

Total number of hits satisfying chosen parameters: 26661

Minimum DB seq length: 0

Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 1500 summaries

Database : Published Applications AA\_US09\*

- 1: /cgn2\_5/ptodata/1/pubpaa/US09\_NEW\_PUB pep:\*\*
- 2: /cgn2\_6/ptodata/1/pubpaa/US06\_NEW\_PUB pep:\*\*
- 3: /cgn2\_6/ptodata/1/pubpaa/US08\_NEW\_PUB pep:\*\*
- 4: /cgn2\_6/ptodata/1/pubpaa/US08\_NEW\_PUB pep:\*\*
- 5: /cgn2\_6/ptodata/1/pubpaa/PCT\_NEW\_PUB pep:\*\*
- 6: /cgn2\_6/ptodata/1/pubpaa/US10\_NEW\_PUB pep:\*\*
- 7: /cgn2\_6/ptodata/1/pubpaa/US11\_NEW\_PUB pep:\*\*
- 8: /cgn2\_6/ptodata/1/pubpaa/US60\_NEW\_PUB pep:\*\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Match	Length	DB ID	Description
1	231.5	15.4	123	6	US-10-821-234-1241
2	176.5	11.7	3002	6	US-10-821-234-916
3	175	11.6	652	6	US-10-821-234-1016
4	165	11.0	1238	7	US-11-078-735-21
5	160.5	10.7	618	7	US-11-137-465-18
6	158.5	10.5	387	7	US-11-137-465-57
7	158	10.5	997	7	US-11-113-424-37
8	156	10.4	999	7	US-11-113-424-36
9	155.5	10.3	961	7	US-11-113-424-35
10	154	10.2	497	6	US-10-821-234-1119
11	153	10.2	1218	7	US-11-078-735-20
12	152.5	10.1	964	7	US-11-137-465-58
13	152.5	10.1	965	7	US-11-113-424-2
14	152.5	10.1	965	7	US-11-147-047-51
15	150	10.0	1671	7	US-11-113-424-38
16	150	10.0	671	7	US-11-029-003-6
17	148	9.8	484	7	US-11-078-735-43
18	148	9.8	723	7	US-11-078-735-17
19	146.5	9.7	545	7	US-11-102-240-110
20	145.5	9.7	332	7	US-11-078-735-33
21	145.5	9.7	332	7	US-11-078-735-51
22	145.5	9.7	369	7	US-11-078-735-38
23	145	9.6	1400	6	US-10-821-234-1045
24	144.5	9.6	696	7	US-11-029-003-8
25	144.5	9.6	685	7	US-11-078-735-19
26	141.5	9.4	897	7	US-11-137-465-35
27	141.5	9.4	897	7	US-11-137-465-35

Sequence 36, Appl	US-11-137-465-36	993	9.4	32	141.5
Sequence 1076, Ap	US-10-821-234-1076	3717	9.3	33	140
Sequence 38, Appl	US-11-147-047-38	383	9.3	35	139.5
Sequence 1310, Ap	US-10-821-234-1310	986	9.3	36	139.5
Sequence 30, Appl	US-11-078-735-30	291	9.2	37	138.5
Sequence 6, Appl	US-11-096-051-6	821	9.2	38	138
Sequence 20, Appl	US-11-096-051-20	862	9.2	39	138
Sequence 4, Appl	US-11-096-051-4	2376	9.2	40	138
Sequence 2, Appl	US-11-096-051-2	2715	9.2	41	138
Sequence 10, Appl	US-11-096-051-10	2721	9.2	42	138
Sequence 8, Appl	US-11-096-051-8	2725	9.2	43	138
Sequence 1412, Ap	US-10-821-234-1412	703	9.1	44	137
Sequence 51, Appl	US-11-113-424-51	2715	9.1	45	137
Sequence 18, Appl	US-11-113-424-18	296	8.9	47	134.5
Sequence 39, Appl	US-11-113-424-39	956	8.7	48	130.5
Sequence 52, Appl	US-11-113-424-52	2725	8.5	49	127.5
Sequence 12, Appl	US-11-096-051-12	257	8.4	50	126
Sequence 1216, Ap	US-10-821-234-1216	143	8.3	51	125
Sequence 16, Appl	US-11-067-121-16	757	8.3	52	125
Sequence 14, Appl	US-11-113-424-14	2769	8.2	53	124
Sequence 53, Appl	US-11-113-424-53	2515	8.1	54	122
Sequence 236, Ap	US-10-821-234-1236	112	8.0	55	121
Sequence 28, Appl	US-11-113-424-28	747	7.7	56	116.5
Sequence 26, Appl	US-11-113-424-26	747	7.7	57	116.5
Sequence 1164, Ap	US-10-821-234-1164	765	7.6	58	115
Sequence 6, Appl	US-11-067-121-6	755	7.4	60	112
Sequence 2, Appl	US-10-945-853-2	188	7.3	61	109.5
Sequence 1155, Ap	US-10-821-234-1155	4419	7.3	62	109.5
Sequence 4, Appl	US-10-831-997-4	961	7.2	64	108.5
Sequence 1, Appl	US-10-945-853-1	188	7.1	65	107.5
Sequence 2, Appl	US-11-085-775-2	3500	7.0	67	106
Sequence 1116, Ap	US-11-108-172-1116	5405	6.4	68	96
Sequence 8, Appl	US-10-478-345-8	321	6.3	69	95
Sequence 86, Appl	US-11-108-528-86	1613	6.2	70	93.5
Sequence 4, Appl	US-10-478-345-4	336	6.2	71	93
Sequence 2, Appl	US-10-478-345-2	354	6.1	72	92.5
Sequence 17, Appl	US-11-067-121-17	1018	6.1	73	92.5
Sequence 110, App	US-10-986-501-110	457	6.0	74	91
Sequence 84, Appl	US-11-108-528-84	1613	6.0	75	91
Sequence 184, App	US-11-113-424-184	126	5.8	76	88
Sequence 2, Appl	US-10-831-997-2	1170	5.8	77	88
Sequence 28, Appl	US-11-046-456-28	1170	5.8	78	88
Sequence 28, Appl	US-11-076-187-2	468	5.8	79	88
Sequence 38, Appl	US-11-102-240-38	720	5.8	80	87.5
Sequence 1068, Ap	US-11-108-172-1068	5179	5.8	82	87
Sequence 1, Appl	US-10-844-035-1	372	5.7	84	86
Sequence 46, Appl	US-10-980-388-46	575	5.7	85	86
Sequence 1063, Ap	US-10-821-234-1063	486	5.7	87	85.5
Sequence 114, App	US-11-092-140-114	411	5.6	88	85
Sequence 110, App	US-11-092-140-110	488	5.6	89	85
Sequence 7, Appl	US-10-478-345-6	357	5.6	90	84.5
Sequence 6, Appl	US-11-067-121-7	1028	5.6	91	84
Sequence 1300, Ap	US-10-821-234-1300	280	5.5	92	83.5
Sequence 40, Appl	US-10-980-388-40	544	5.5	93	83
Sequence 5, Appl	US-11-076-187-5	833	5.5	94	83
Sequence 30, Appl	US-10-967-527A-30	308	5.5	95	82.5
Sequence 70, Appl	US-11-113-424-70	620	5.5	96	82.5
Sequence 2, Appl	US-11-139-499-2	234	5.4	97	82
Sequence 1532, Ap	US-10-821-234-1532	705	5.4	98	82
Sequence 82, Appl	US-11-108-528-82	1614	5.4	99	81.5
Sequence 80, Appl	US-11-108-528-80	1615	5.4	100	81.5
Sequence 7, Appl	US-11-021-441-7	502	5.4	101	81
Sequence 9, Appl	US-11-021-441-9	563	5.4	102	81
Sequence 15, Appl	US-11-021-441-15	574	5.4	103	81
Sequence 11, Appl	US-11-021-441-11	581	5.4	104	81
Sequence 13, Appl	US-11-021-441-13	581	5.4	105	81
Sequence 4, Appl	US-11-021-441-4	1035	5.4	106	81
Sequence 1539, Ap	US-10-821-234-1539	165	5.3	107	80.5
Sequence 4, Appl	US-11-153-880-4	350	5.3	108	80.5
Sequence 2, Appl	US-11-064-769-2	419	5.3	109	80.5
Sequence 1357, Ap	US-11-153-880-2	419	5.3	110	80.5
	US-11-821-234-1357	405	5.3	111	80

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OM protein - protein search, using sw model

Run on: December 3, 2005, 23:21:27 ; Search time 66 Seconds  
(without alignments)  
1728.294 Million cell updates/sec

Perfect score: 1505  
Sequence: 1 MRGSOVLLMLLAVGVT.....SRQISFLERQLGSCCKKDS 273

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 1867569 seqs, 417829326 residues

Total number of hits satisfying chosen parameters: 1867569

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 1500 summaries

- Database :
- 1: /cgn2\_6/ptodata/1/pubpaa/US07\_PUBCOMB.pep.\*
  - 2: /cgn2\_6/ptodata/1/pubpaa/US08\_PUBCOMB.pep.\*
  - 3: /cgn2\_6/ptodata/1/pubpaa/US09\_PUBCOMB.pep.\*
  - 4: /cgn2\_6/ptodata/1/pubpaa/US10A\_PUBCOMB.pep.\*
  - 5: /cgn2\_6/ptodata/1/pubpaa/US10B\_PUBCOMB.pep.\*
  - 6: /cgn2\_6/ptodata/1/pubpaa/US11\_PUBCOMB.pep.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	1505	100.0	273	3	US-09-790-264-10
75	1505	100.0	273	4	US-10-167-749-506
76	1505	100.0	273	4	US-10-167-749-508
125	1505	100.0	273	4	US-10-219-065-178
151	1505	100.0	273	4	US-10-269-353-10
241	1505	100.0	273	4	US-10-170-481A-506
242	1505	100.0	273	4	US-10-170-481A-508
245	1505	100.0	273	4	US-10-210-028-506
246	1505	100.0	273	4	US-10-210-028-508
255	1505	100.0	273	4	US-10-162-521A-506
256	1505	100.0	273	4	US-10-162-521A-508
281	1505	100.0	273	5	US-10-287-971-102
286	1505	100.0	273	5	US-10-918-851-506
287	1505	100.0	273	5	US-10-918-851-508
288	1505	100.0	273	5	US-10-900-926-10
289	1505	100.0	273	5	US-10-805-667-506
290	1505	100.0	273	5	US-10-805-667-508
291	1505	100.0	273	5	US-10-897-359-506
292	1505	100.0	273	5	US-10-897-359-508
293	1505	100.0	273	5	US-10-893-802-506
294	1505	100.0	273	5	US-10-893-802-508
295	1505	100.0	273	5	US-10-897-360-506
296	1505	100.0	273	5	US-10-897-360-508
299	1505	100.0	273	6	US-11-129-762-506
300	1505	100.0	273	6	US-11-129-762-508
325	1504	99.9	273	3	US-09-852-472-2
326	1504	99.9	273	4	US-10-167-749-510
372	1504	99.9	273	4	US-10-170-481A-510

Sequence 510, App	273	4	US-10-210-028-510
Sequence 510, App	273	4	US-10-162-521A-510
Sequence 84, Appl	273	4	US-10-287-971-84
Sequence 100, App	273	4	US-10-287-971-100
Sequence 900, App	273	4	US-10-408-765A-900
Sequence 510, App	273	5	US-10-918-851-510
Sequence 510, App	273	5	US-10-805-667-510
Sequence 510, App	273	5	US-10-897-359-510
Sequence 510, App	273	5	US-10-893-802-510
Sequence 510, App	273	5	US-10-897-360-510
Sequence 510, App	273	6	US-11-129-762-510
Sequence 88, Appl	280	4	US-10-287-971-88
Sequence 92, Appl	285	4	US-10-425-114-37265
Sequence 37265, A	293	4	US-10-287-971-96
Sequence 96, Appl	254	3	US-09-852-472-3
Sequence 3, Appl	251	3	US-09-790-264-12
Sequence 12, Appl	251	4	US-10-269-353-12
Sequence 12, Appl	251	5	US-10-900-926-12
Sequence 90, Appl	256	4	US-10-287-971-90
Sequence 94, Appl	256	4	US-10-287-971-94
Sequence 2, Appl	295	4	US-10-167-749-2
Sequence 374, App	295	4	US-10-223-085-374
Sequence 374, App	295	4	US-10-223-084-374
Sequence 374, App	295	4	US-10-223-088-374
Sequence 374, App	295	4	US-10-223-087-374
Sequence 374, App	295	4	US-10-223-083-374
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Sequence 374, App	295	4	US-10-223-082-374
Sequence 2, Appl	295	4	US-10-170-481A-2
Sequence 2, Appl	295	4	US-10-210-028-2
Sequence 2, Appl	295	4	US-10-162-521A-2
Sequence 374, App	295	4	US-10-305-654-374
Sequence 374, App	295	4	US-10-081-056-374
Sequence 2, Appl	295	5	US-10-918-851-2
Sequence 2, Appl	295	5	US-10-805-667-2
Sequence 2, Appl	295	5	US-10-897-359-2
Sequence 2, Appl	295	5	US-10-893-802-2
Sequence 2, Appl	295	6	US-11-129-762-2
Sequence 18, Appl	234	4	US-10-098-871-18
Sequence 15, Appl	275	3	US-09-790-264-15
Sequence 15, Appl	275	4	US-10-269-353-15
Sequence 15, Appl	275	5	US-10-900-926-15
Sequence 19, Appl	258	3	US-09-852-472-19
Sequence 24, Appl	275	3	US-09-852-472-24
Sequence 692, App	255	3	US-09-865-050A-692
Sequence 86, Appl	249	4	US-10-287-971-86
Sequence 218, App	287	3	US-09-764-898-218
Sequence 111, App	288	3	US-09-764-893-111
Sequence 107, App	288	3	US-09-764-881-107
Sequence 841, App	288	3	US-09-764-853-841
Sequence 290, App	288	3	US-09-764-888-290
Sequence 107, App	288	4	US-10-073-865-107
Sequence 111, App	288	4	US-10-242-747-107
Sequence 670, App	314	3	US-09-764-853-670
Sequence 15, Appl	169	3	US-09-852-472-15
Sequence 13, Appl	158	3	US-09-852-472-13
Sequence 16, Appl	181	3	US-09-852-472-16
Sequence 20, Appl	247	4	US-10-269-353-20
Sequence 20, Appl	247	4	US-10-269-353-20
Sequence 14, Appl	247	5	US-10-900-926-14
Sequence 17638, A	293	3	US-09-852-472-17
Sequence 77, Appl	509	6	US-11-097-143-16638
Sequence 77, Appl	1574	3	US-09-825-751A-77
Sequence 77, Appl	1574	5	US-10-851-438-77
Sequence 6, Appl	553	3	US-09-981-649A-6
Sequence 6, Appl	553	4	US-10-124-986-6
Sequence 6, Appl	553	4	US-10-136-227A-6

W. Harvey, L.  
09/19/78/191  
Seq ID 506  
Sub 7 Issues dls

GenCore version 5.1.6  
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OM protein - protein search, using sw model

Run on: December 3, 2005, 23:18:21 ; Search time 24 Seconds

(without alignments)  
940.436 Million cell updates/sec

Perfect score: 1505

Sequence: 1 MRGQEVLLMVLVAVGGT.....SEQISFLEQLGSCCKKDS 273

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 572060 seqs, 82675679 residues

Total number of hits satisfying chosen parameters: 572060

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 1500 summaries

Database :

- 1: /cgn2\_6/ptodata/1/iaa/5 COMB.pep.\*
- 2: /cgn2\_6/ptodata/1/iaa/6 COMB.pep.\*
- 3: /cgn2\_6/ptodata/1/iaa/H COMB.pep.\*
- 4: /cgn2\_6/ptodata/1/iaa/PCRUS COMB.pep.\*
- 5: /cgn2\_6/ptodata/1/iaa/RE COMB.pep.\*
- 6: /cgn2\_6/ptodata/1/iaa/backfiles1.pep.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match %	Length	ID	Description
9	1180	78.4	278	2	US-09-724-864-52
10	254.5	16.9	553	2	US-09-249-697A-6
11	254.5	16.9	553	2	US-09-363-316B-6
12	254.5	16.9	553	2	US-10-136-227A-6
13	254.5	16.9	553	2	US-09-981-649A-6
14	252.5	16.8	284	2	US-09-312-283C-389
17	251.5	16.7	553	2	US-09-249-697A-19
18	251.5	16.7	553	2	US-09-363-316B-24
19	251.5	16.7	553	2	US-10-136-227A-24
20	251.5	16.7	553	2	US-09-981-649A-24
21	251.5	16.7	554	2	US-10-136-227A-32
22	251.5	16.7	554	2	US-09-981-649A-32
23	251.5	16.7	594	2	US-09-949-016-8175
24	249.5	16.6	554	2	US-10-136-227A-30
25	249.5	16.6	554	2	US-09-981-649A-30
26	249.5	16.6	559	2	US-10-136-227A-28
27	249.5	16.6	559	2	US-09-981-649A-28
41	216.5	14.4	502	2	US-09-363-316B-18
42	216.5	14.4	502	2	US-10-136-227A-18
43	216.5	14.4	502	2	US-09-981-649A-18
44	216.5	14.4	537	2	US-09-249-697A-4
45	216.5	14.4	537	2	US-09-363-316B-4
46	216.5	14.4	537	2	US-10-136-227A-4
47	216.5	14.4	537	2	US-09-981-649A-4
48	215.5	14.3	100	2	US-09-249-697A-3
49	215.5	14.3	100	2	US-09-363-316B-3
50	215.5	14.3	100	2	US-10-136-227A-3
51	215.5	14.3	100	2	US-09-981-649A-3

1964	2	US-09-467-997-1	13.0	196	52	Sequence 1, Appli
678	1	US-08-282-141-2	12.6	189	53	Sequence 2, Appli
678	1	US-08-435-434-2	12.6	189	54	Sequence 2, Appli
678	1	US-08-435-436-2	12.6	189	55	Sequence 2, Appli
678	1	US-08-438-863-2	12.6	189	56	Sequence 2, Appli
678	1	US-08-438-864-2	12.6	189	57	Sequence 2, Appli
678	1	US-08-438-862-2	12.6	189	58	Sequence 2, Appli
678	2	US-08-628-747-2	12.6	189	59	Sequence 2, Appli
678	2	US-08-402-253-2	12.6	189	60	Sequence 2, Appli
678	2	US-08-443-866B-2	12.6	189	61	Sequence 2, Appli
673	1	US-08-282-141-3	12.2	184	62	Sequence 3, Appli
673	1	US-08-435-434-1	12.2	184	63	Sequence 1, Appli
673	1	US-08-435-436-1	12.2	184	64	Sequence 1, Appli
673	1	US-08-438-863-1	12.2	184	65	Sequence 1, Appli
673	1	US-08-438-864-1	12.2	184	66	Sequence 1, Appli
673	2	US-08-438-862-1	12.2	184	67	Sequence 1, Appli
673	2	US-08-628-747-1	12.2	184	68	Sequence 1, Appli
673	2	US-08-402-253-1	12.2	184	69	Sequence 1, Appli
673	2	US-08-443-866B-1	12.2	184	70	Sequence 1, Appli
816	1	US-08-820-170A-37	12.1	182.5	71	Sequence 37, Appl
816	2	US-09-055-699-37	12.1	182.5	72	Sequence 37, Appl
816	2	US-09-273-565-37	12.1	182.5	73	Sequence 37, Appl
816	2	US-09-565-538-37	12.1	182.5	74	Sequence 37, Appl
816	2	US-09-661-468-37	12.1	182.5	75	Sequence 37, Appl
816	2	US-09-976-165-37	12.1	182.5	76	Sequence 37, Appl
810	1	US-08-820-170A-34	12.1	182	77	Sequence 34, Appl
810	2	US-09-055-699-34	12.1	182	78	Sequence 34, Appl
810	2	US-09-273-565-34	12.1	182	79	Sequence 34, Appl
810	2	US-09-565-538-34	12.1	182	80	Sequence 34, Appl
810	2	US-09-661-468-34	12.1	182	81	Sequence 34, Appl
810	2	US-09-976-165-34	12.1	182	82	Sequence 34, Appl
1935	2	US-09-949-016-10403	11.7	176.5	83	Sequence 10403, A
2871	2	US-09-538-092-1076	11.7	176.5	84	Sequence 1076, Ap
1581	2	US-09-949-002-414	11.7	176	85	Sequence 414, App
2321	2	US-09-230-652-2	11.7	175.5	86	Sequence 2, Appli
652	1	US-08-751-305-2	11.6	175	87	Sequence 2, Appli
1587	2	US-09-949-002-354	11.6	175	88	Sequence 354, App
2703	1	US-08-185-432-19	11.6	175	89	Sequence 19, Appl
2703	2	US-08-899-232-4	11.6	175	90	Sequence 4, Appli
2703	2	US-09-121-457-4	11.6	175	91	Sequence 4, Appli
1246	2	US-09-919-497-85	11.5	173.5	92	Sequence 85, Appl
4247	2	US-09-961-403-14	11.5	173.5	93	Sequence 14, Appl
4588	2	US-10-025-225-8	11.5	173	94	Sequence 8, Appli
1833	2	US-08-479-722B-2	11.5	172.5	95	Sequence 2, Appli
1833	2	US-09-592-685-2	11.5	172.5	96	Sequence 2, Appli
1833	4	FCT-US95-02251-18	11.5	172.5	97	Sequence 18, Appl
830	2	US-08-872-855-11	11.3	170	98	Sequence 11, Appl
4585	2	US-10-025-225-6	11.3	169.5	99	Sequence 6, Appli
4589	2	US-10-025-225-6	11.3	169.5	100	Sequence 2, Appli
833	1	US-08-864-534-6	11.2	169	101	Sequence 6, Appli
833	1	US-08-083-590A-2	11.2	169	102	Sequence 2, Appli
833	1	US-08-465-500-6	11.2	169	103	Sequence 6, Appli
833	1	US-08-346-126-6	11.2	169	104	Sequence 6, Appli
833	1	US-08-346-128-6	11.2	169	105	Sequence 6, Appli
833	2	US-08-532-384-2	11.2	169	106	Sequence 2, Appli
833	2	US-08-893-828-6	11.2	169	107	Sequence 2, Appli
833	2	US-09-310-685-2	11.2	169	108	Sequence 2, Appli
1055	2	US-09-214-278-2	11.2	168	109	Sequence 2, Appli
1055	1	US-08-400-159-10	11.2	168	110	Sequence 10, Appl
1193	2	US-08-611-729A-10	11.2	168	111	Sequence 10, Appl
1193	2	US-09-195-524-10	11.2	168	112	Sequence 10, Appl
1193	2	US-09-310-685-8	11.2	168	113	Sequence 8, Appli
832	2	US-08-981-392-6	11.0	166	114	Sequence 6, Appli
832	2	US-09-908-322-6	11.0	166	115	Sequence 8, Appli
1065	1	US-08-400-159-8	11.0	165	116	Sequence 3, Appli
1212	2	US-09-214-278-3	11.0	165	117	Sequence 3, Appli
1212	2	US-09-855-722-3	11.0	165	118	Sequence 3, Appli
1238	2	US-09-214-278-5	11.0	165	119	Sequence 5, Appli
1238	2	US-09-855-722-5	11.0	165	120	Sequence 5, Appli
1257	2	US-08-611-729A-8	11.0	165	121	Sequence 8, Appli
1257	2	US-09-195-524-8	11.0	165	122	Sequence 8, Appli
1257	2	US-09-310-685-6	11.0	165	123	Sequence 6, Appli
1257	2	US-09-310-685-6	11.0	165	124	Sequence 6, Appli